



SEQUENCE LISTING

<110> UENAI, TSUJI, OKAMURA, HIRO
KAMBARA, HIDEAKI
OKANO, KAZUNORI

<120> METHOD FOR ASSAYING DNA FRAGMENTS IN MIXTURE

<130> H-853-02

<140> 09/842,791

<141> 2001-04-27

<150> 09/413,814

<151> 1999-10-07

<150> JP 10-286286

<151> 1998-10-08

<160> 4

<170> PatentIn Ver. 2.1

<210> 1

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<220>

<221> modified_base

<222> (21)..(22)

<223> a, t, c, or g

<400> 1

tcatcacctt ctccacgac nn

22

<210> 2

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<220>

<221> modified_base

<222> (21)..(22)

<223> a, t, c, or g

<400> 2

ttctccactc atcaccgac nn

22

<210> 3
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<220>
<221> modified_base
<222> (21)..(22)
<223> a, t, c, or g

<400> 3
acaagacgat cagcaggatc nn

22

<210> 4
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<220>
<221> modified_base
<222> (21)..(22)
<223> a, t, c, or g

<400> 4
acaagcagat cagacggatc nn

22